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REMARKS

It is believed that the above amendments bring the application in compliance with 37 CFR 1.821-1.825.

In view of the above amendments, reconsideration and allowance of the above-identified application is respectfully requested.

Attached hereto is a marked-up version of the changes made to the specification by the current amendment. The attached page is captioned "Version with markings to show changes made."

Respectfully submitted, ...

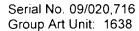
Mid ad and Himchel

Marianne H. Michel Attorney for Applicant

Registration No. 35,286

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VERSION WITH MARKINGS TO SHOW CHANGES MADE

In showing the changes, deleted material is shown as a strike through, and inserted material is shown as underlined.

In the Specification:

Paragraph beginning at line 4 of page 28 has been amended as follows:

	Primers	are	designated	as	HTPCR1	Seq.	8	(5'-
AGTA	TAAGTA	AACACA	ACCATCACACCC	TTGA	GGCCCTTGC	TGGTGG	CCAT	GGT
G-3')		and	HTPCR2		Seq.	9		(5'-
ССТС	CACATCC	CTTAG	GCCTAAGTTCG.	ACGT	CGGGCCCTC	CTAGTCG	ACGG	SATC
CA-3'). These	primers :	are used in a PCR	react	tion to amplify	alpha hor	dothior	nin by
conve	entional m	ethods.	The resulting PC	R pro	duct is purifie	d and sul	bclone	d into
the B	amHI/Nco	l digeste	ed pBSKP vector (Strata	agene, LaJolla	, CA) and	l seque	enced
on bo	th strands	to confi	rm its identity. Th	e clon	ie is designate	d pBSKP	-HT (se	∍q. ID
1). F	Primers ar	e desigr	ned for single stra	anded	DNA site-dire	ected mut	agene	sis to
introd	uce 12 co	odons fo	r lysine, based on	the p	peptide structu	re of hore	dothion	ıin 12
(Ref:	Rao et a	<i>l</i> . 1994	<u>Protein Engineerir</u>	ng 7(1	12):1485-1493)	and are	desig	nated
HT12	mut1 <u>Sec</u>	<u>ı. 10</u> (5'	-AGCGGAAAATG	ccc	SAAAGGCTTC	CCCAAA	\TTGG	C-3'),
HT12	mut2		Seq.		11			(5'-
TGCGCAGGCGTCTGCAAGTGTAAGCTGACTAGTAGCGGAAAATGC-3'),								
HT12	mut3		<u>Seq.</u>		12			(5'-
TACA	ACCTTT	GCAAAG	STCAAAGGCGCC	AAGA	AGCTTTGCG	CAGGC	STCTG	3 -3'),
HT12	mut4		Seq.		13			(5'-
GCAA	GAGTTG	CTGCA	AGAGTACCCTGC	GAA	GGAAGTGCT	ACAACC	TTTGC	:-3')

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Table 2 on page 40 has been amended as follows:

Table 2: SEQUENCE INFORMATION

Table 2. SEQUENCE IN ORMATION									
SEQUENCE ID	PROMOTER	GENE							
Seq. 1: pBSKP-HT	None	3361-2947							
Seq. 2: pBSKP-HT12	None	3361-2947							
Seq. 3: PHP8001gz::HT12::gz expression vector	676-2198	2199-2612							
Seq. 4: PHP7999 glb1::HT12::glb1 expression vector	3271-1834	1834-1420							
Seq. 5: PHP5025 wx::HT::wx expression vector	43-1342	1343-1757							
Seq. 6: PHP 11260 gz::ESA::gz expression vector	676-2198	2199-2675							
Seq. 7: PHP11427 gz::BHL::gz	676-2198	2199-2450							
Seq. 8-13: artificial sequence primers									
Seq. 14: Pea albumin, nucleotide sequence									
Seq. 15: Pea albumin, protein sequence									

<u>Seq. 16:</u> sulfur-rich 15KD maize protein, nucleotide sequence

<u>Seq. 17:</u> sulfur-rich 15KD maize protein, protein sequence

<u>Seq. 18: methionine-rich 10KD maize protein, nucleotide sequence</u>

Seq. 19: methionine-rich 10KD maize protein, protein sequence

Seq. 20: sulfur-rich rice prolamine, nucleotide sequence

Seq. 21: sulfur-rich rice prolamine, protein sequence

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Seq. 22: wheat endosperm purothionin, protein sequence